

# SEQUENCE LISTING

<110> RUSCH, Douglas et al

<120> ISOLATED HUMAN KINASE PROTEINS, NUCLEIC  
ACID MOLECULES ENCODING HUMAN KINASE PROTEINS, AND USES  
THEREOF

<130> CL001099-CIP-DIV2

<140> To be assigned

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<141> 2002-10-22

<150> 09/849,334

<151> 2001-05-07

<150> 09/773,371

<151> 2001-02-01

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<212> DNA

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Ile Asp Lys Glu Leu Gln Met Arg Thr Gly Ala Glu Asn Leu Tyr Arg
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Ser Tyr Glu Ala Glu Ile Arg Glu Leu Glu Ala Leu Arg Gln Ala Met
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715

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<400> 4
Met Ile Leu Glu Glu Arg Pro Asp Gly Gln Gly Thr Gly Glu Glu Ser
 1             5             10             15
Ser Arg Pro Gln Asp Asp Gly Ser Ile Arg Lys Gly Tyr Gly Ser Phe
      20             25             30
Val Gln Asn Gln Pro Gly Gln Leu Ser His Arg Ala Arg Leu His
    35             40             45

```

Gln	Gln	Ile	Ser	Lys	Glu	Leu	Arg	Met	Arg	Thr	Gly	Ala	Glu	Asn	Leu	
50						55					60					
Tyr	Arg	Ala	Thr	Ser	Asn	Thr	Trp	Val	Arg	Glu	Thr	Val	Ala	Leu	Glu	
65					70					75					80	
Leu	Ser	Tyr	Val	Asn	Ser	Asn	Leu	Gln	Leu	Leu	Lys	Glu	Glu	Leu	Ala	
				85					90					95		
Glu	Leu	Ser	Thr	Ser	Val	Asp	Val	Asp	Gln	Pro	Glu	Gly	Glu	Gly	Ile	
			100					105					110			
Thr	Ile	Pro	Met	Ile	Pro	Leu	Gly	Leu	Lys	Glu	Thr	Lys	Glu	Leu	Asp	
		115					120					125				
Trp	Ala	Thr	Pro	Leu	Lys	Glu	Leu	Ile	Ser	Glu	His	Phe	Gly	Glu	Asp	
130						135					140					
Gly	Thr	Ser	Phe	Glu	Thr	Glu	Ile	Gln	Glu	Leu	Glu	Asp	Leu	Arg	Gln	
145				150					155					160		
Ala	Thr	Arg	Thr	Pro	Ser	Arg	Asp	Glu	Ala	Gly	Leu	Asp	Leu	Leu	Ala	
				165				170					175			
Ala	Tyr	Tyr	Ser	Gln	Leu	Cys	Phe	Leu	Asp	Ala	Arg	Phe	Phe	Ser	Pro	
			180					185				190				
Ser	Arg	Ser	Pro	Gly	Leu	Leu	Phe	His	Trp	Tyr	Asp	Ser	Leu	Thr	Gly	
		195					200				205					
Val	Pro	Ala	Gln	Gln	Arg	Ala	Leu	Ala	Phe	Glu	Lys	Gly	Ser	Val	Leu	
210						215					220					
Phe	Asn	Ile	Gly	Ala	Leu	His	Thr	Gln	Ile	Gly	Ala	Arg	Gln	Asp	Cys	
225				230					235					240		
Ser	Cys	Thr	Glu	Gly	Thr	Asn	His	Ala	Ala	Glu	Ala	Phe	Gln	Arg	Ala	
				245				250					255			
Ala	Gly	Ala	Phe	Arg	Leu	Leu	Arg	Glu	Asn	Phe	Ser	His	Ala	Pro	Ser	
			260					265				270				
Pro	Asp	Met	Ser	Ala	Ala	Ser	Leu	Ser	Met	Leu	Glu	Gln	Leu	Met	Ile	
		275				280					285					
Ala	Gln	Ala	Gln	Glu	Cys	Ile	Phe	Lys	Gly	Leu	Leu	Leu	Pro	Ala	Ser	
290						295					300					
Ala	Thr	Pro	Asp	Ile	Cys	Pro	Asp	Gln	Leu	Gln	Leu	Ala	Gln	Glu	Ala	
305				310				315						320		
Ala	Gln	Val	Ala	Thr	Glu	Tyr	Gly	Leu	Val	His	Arg	Ala	Met	Ala	Gln	
				325				330					335			
Pro	Pro	Val	Arg	Asp	Tyr	Leu	Pro	Ala	Ser	Trp	Thr	Asn	Leu	Ala	His	
			340					345				350				
Val	Lys	Ala	Glu	His	Phe	Cys	Ala	Leu	Ala	His	Tyr	His	Ala	Ala	Met	
		355				360					365					
Ala	Leu	Cys	Glu	Ser	His	Pro	Ala	Lys	Gly	Glu	Leu	Ala	Arg	Gln	Glu	
370						375				380						
His	Val	Phe	Gln	Pro	Ser	Thr	Pro	His	Glu	Pro	Leu	Gly	Pro	Thr	Leu	
385				390					395					400		
Pro	Gln	His	Pro	Glu	Asp	Arg	Arg	Lys	Leu	Ala	Lys	Ala	His	Leu	Lys	
				405				410					415			
Arg	Ala	Ile	Leu	Gly	Gln	Glu	Glu	Ala	Leu	Arg	Leu	His	Thr	Leu	Cys	
			420					425				430				
Arg	Val	Leu	Arg	Lys	Val	Asp	Leu	Leu	Gln	Val	Val	Val	Thr	Gln	Ala	
		435				440					445					
Leu	Arg	Arg	Ser	Leu	Ala	Lys	Tyr	Ser	Gln	Leu	Glu	Arg	Glu	Asp	Asp	
450						455				460						
Phe	Phe	Glu	Ala	Thr	Glu	Ala	Pro	Asp	Ile	Gln	Pro	Lys	Thr	His	Gln	
465				470					475					480		
Thr	Pro	Glu	Gly	Pro	Leu	Ser	Val	Phe	Ser	Thr	Lys	Asn	Arg	Trp	Gln	
				485				490				495				
Leu	Val	Gly	Pro	Val	His	Met	Thr	Arg	Gly	Glu	Gly	Gly	Phe	Gly	Phe	

			500					505					510				
Thr	Leu	Arg	Gly	Asp	Ser	Pro	Val	Leu	Ile	Ala	Ala	Val	Val	Pro	Gly		
		515					520					525					
Gly	Gln	Ala	Glu	Ser	Ala	Gly	Leu	Lys	Glu	Gly	Asp	Tyr	Ile	Val	Ser		
	530					535					540						
Val	Asn	Gly	Gln	Pro	Cys	Lys	Trp	Trp	Lys	His	Leu	Glu	Val	Val	Thr		
545					550					555					560		
Gln	Leu	Arg	Ser	Met	Gly	Glu	Glu	Gly	Val	Ser	Leu	Gln	Val	Val	Ser		
			565					570						575			
Leu	Leu	Pro	Ser	Pro	Glu	Pro	Arg	Gly	Thr	Gly	Pro	Arg	Arg	Ala	Ala		
		580						585					590				
Leu	Leu	Trp	Asn	Gln	Arg	Glu	Cys	Gly	Phe	Glu	Thr	Pro	Met	Pro	Thr		
	595						600					605					
Arg	Thr	Arg	Pro	Trp	Pro	Ile	Leu	Gly	Trp	Ser	Arg	Lys	Asn	Lys	Gln		
	610					615					620						
Gly	Lys	Thr	Gly	Ser	His	Pro	Asp	Pro	Cys								
625					630												